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Toyohide SHINKAWA
Naoko YAMANE
Motoo YAMASAKI
Nobuo HANAI

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 aacactgcat cctgatgcct ctgcacactt ccattcttta gatgacatct actattttgg 780
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 aatccccatg gaacctggag atatcattgg tgtggctgga aaccattgga atggttactc 900
 taaagggtgc aacagaaaac taggaaaaac aggcctgtac cttcctaca aagtccgaga 960

gaagatagaa acggtcaag

979

<210> 7

<211> 979

<212> DNA

<213> Rattus norvegicus

<400> 7

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<210> 8

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 8

aagtataagc ttacatggat gacgatatcg ctgcgtcgt 40

<210> 9

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 9

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<210> 10
<211> 40
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<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 10
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<210> 11
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<400> 11
ctccaattat gaatttatta gtg 23

<210> 12
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<400> 12
ggatgtttga agccaagctt cttgg 25

<210> 13
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<220>
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<400> 13
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<210> 14
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<220>
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<400> 14
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<210> 15
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<400> 15
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<210> 16
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<210> 17
<211> 24
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<400> 17
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<210> 18
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<212> DNA
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<400> 18
caggaaggaa ggctggaaga gacg 24

<210> 19
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<212> DNA
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<220>
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<400> 19
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<210> 20
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<400> 20
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<210> 21
<211> 24
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 21

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24

<210> 22

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 22

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23

<210> 23

<211> 575

<212> PRT

<213> Cricetulus griseus

<400> 23

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Asn Asp His Pro Asp His Ser Ser Arg Glu Leu Ser Lys Ile Leu Ala
35 40 45

Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala
50 55 60

Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr
65 70 75 80

Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln
85 90 95

Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Asp Leu Gly Lys Asp His
100 105 110

Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe
115 120 125

Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys Lys Leu Glu Gly Asn Glu
130 135 140

Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu
145 150 155 160

Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala
165 170 175

Gly Glu Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln
180 185 190

Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg
195 200 205

Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu
210 215 220

His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr
 225 230 235 240
 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu
 245 250 255
 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu
 260 265 270
 Ser Thr Gly His Trp Ser Gly Glu Val Lys Asp Lys Asn Val Gln Val
 275 280 285
 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu
 290 295 300
 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His
 305 310 315 320
 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile
 325 330 335
 Arg Pro Gln Pro Trp Leu Glu Arg Glu Ile Glu Glu Thr Thr Lys Lys
 340 345 350
 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp
 355 360 365
 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val
 370 375 380
 His Val Glu Glu His Phe Gln Leu Leu Glu Arg Arg Met Lys Val Asp
 385 390 395 400
 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Ser Leu Leu Lys Glu
 405 410 415
 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile
 420 425 430
 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg
 435 440 445
 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val
 450 455 460
 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln
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 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile
 485 490 495
 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro
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 His Gln Pro Arg Thr Lys Glu Glu Ile Pro Met Glu Pro Gly Asp Ile
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 Ile Gly Val Ala Gly Asn His Trp Asn Gly Tyr Ser Lys Gly Val Asn
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<210> 24
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 <212> PRT
 <213> Mus musculus

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 35 40 45
 Lys Leu Glu Arg Leu Lys Gln Gln Asn Glu Asp Leu Arg Arg Met Ala
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 Glu Ser Leu Arg Ile Pro Glu Gly Pro Ile Asp Gln Gly Thr Ala Thr
 65 70 75 80
 Gly Arg Val Arg Val Leu Glu Glu Gln Leu Val Lys Ala Lys Glu Gln
 85 90 95
 Ile Glu Asn Tyr Lys Lys Gln Ala Arg Asn Gly Leu Gly Lys Asp His
 100 105 110
 Glu Ile Leu Arg Arg Arg Ile Glu Asn Gly Ala Lys Glu Leu Trp Phe
 115 120 125
 Phe Leu Gln Ser Glu Leu Lys Lys Leu Lys His Leu Glu Gly Asn Glu
 130 135 140
 Leu Gln Arg His Ala Asp Glu Ile Leu Leu Asp Leu Gly His His Glu
 145 150 155 160
 Arg Ser Ile Met Thr Asp Leu Tyr Tyr Leu Ser Gln Thr Asp Gly Ala
 165 170 175
 Gly Asp Trp Arg Glu Lys Glu Ala Lys Asp Leu Thr Glu Leu Val Gln
 180 185 190
 Arg Arg Ile Thr Tyr Leu Gln Asn Pro Lys Asp Cys Ser Lys Ala Arg
 195 200 205
 Lys Leu Val Cys Asn Ile Asn Lys Gly Cys Gly Tyr Gly Cys Gln Leu
 210 215 220
 His His Val Val Tyr Cys Phe Met Ile Ala Tyr Gly Thr Gln Arg Thr
 225 230 235 240
 Leu Ile Leu Glu Ser Gln Asn Trp Arg Tyr Ala Thr Gly Gly Trp Glu
 245 250 255
 Thr Val Phe Arg Pro Val Ser Glu Thr Cys Thr Asp Arg Ser Gly Leu
 260 265 270
 Ser Thr Gly His Trp Ser Gly Glu Val Asn Asp Lys Asn Ile Gln Val
 275 280 285
 Val Glu Leu Pro Ile Val Asp Ser Leu His Pro Arg Pro Pro Tyr Leu
 290 295 300
 Pro Leu Ala Val Pro Glu Asp Leu Ala Asp Arg Leu Leu Arg Val His
 305 310 315 320
 Gly Asp Pro Ala Val Trp Trp Val Ser Gln Phe Val Lys Tyr Leu Ile
 325 330 335

Arg Pro Gln Pro Trp Leu Glu Lys Glu Ile Glu Glu Ala Thr Lys Lys
 340 345 350
 Leu Gly Phe Lys His Pro Val Ile Gly Val His Val Arg Arg Thr Asp
 355 360 365
 Lys Val Gly Thr Glu Ala Ala Phe His Pro Ile Glu Glu Tyr Met Val
 370 375 380
 His Val Glu Glu His Phe Gln Leu Leu Ala Arg Arg Met Gln Val Asp
 385 390 395 400
 Lys Lys Arg Val Tyr Leu Ala Thr Asp Asp Pro Thr Leu Leu Lys Glu
 405 410 415
 Ala Lys Thr Lys Tyr Ser Asn Tyr Glu Phe Ile Ser Asp Asn Ser Ile
 420 425 430
 Ser Trp Ser Ala Gly Leu His Asn Arg Tyr Thr Glu Asn Ser Leu Arg
 435 440 445
 Gly Val Ile Leu Asp Ile His Phe Leu Ser Gln Ala Asp Phe Leu Val
 450 455 460
 Cys Thr Phe Ser Ser Gln Val Cys Arg Val Ala Tyr Glu Ile Met Gln
 465 470 475 480
 Thr Leu His Pro Asp Ala Ser Ala Asn Phe His Ser Leu Asp Asp Ile
 485 490 495
 Tyr Tyr Phe Gly Gly Gln Asn Ala His Asn Gln Ile Ala Val Tyr Pro
 500 505 510
 His Lys Pro Arg Thr Glu Glu Glu Ile Pro Met Glu Pro Gly Asp Ile
 515 520 525
 Ile Gly Val Ala Gly Asn His Trp Asp Gly Tyr Ser Lys Gly Ile Asn
 530 535 540
 Arg Lys Leu Gly Lys Thr Gly Leu Tyr Pro Ser Tyr Lys Val Arg Glu
 545 550 555 560
 Lys Ile Glu Thr Val Lys Tyr Pro Thr Tyr Pro Glu Ala Glu Lys
 565 570 575

<210> 25
 <211> 18
 <212> PRT
 <213> Homo sapiens

<400> 25
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 1 5 10 15

Pro Cys

<210> 26
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic DNA

<400> 26

cttgtgtgac tcttaactct cagag

25

<210> 27

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 27

ccctcgagat aacttcgtat agc

23

<210> 28

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 28

ggtaggcctc actaactg

18

<210> 29

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 29

catagaaaca agtaacaaca gccag

25

<210> 30

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 30

gagacttcag cccacttcaa ttattggc

28

<210> 31

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 31

gaggccactt gtgtagcgcc aagtg

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<210> 32

<211> 24

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic DNA

<400> 32

aggaagggtgg cgctcatcac gggc

24

<210> 33

<211> 26

<212> DNA

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<223> Description of Artificial Sequence: Synthetic DNA

<400> 33

taaggccaca agtcttaatt gcatcc

26

<210> 34

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic DNA

<400> 34

caggggtgtt cccttgagga ggtggaa

27

<210> 35

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 35

cccctcacgc atgaagcctg gag

23

<210> 36

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 36

ggcaggagac cacctgcga gtgccac

28

<210> 37

<211> 28

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic DNA

<400> 37

ggcgctggct tacccggaga ggaatggg

28

<210> 38

<211> 28
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<223> Description of Artificial Sequence: Synthetic DNA

<400> 38
aaaaggcctc agttagtga ctgtatgg 28

<210> 39
<211> 29
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 39
cgcggtacct caagcgttg ggttggtcc 29

<210> 40
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 40
cccaagcttg ccaccatggc tcacgctccc gctagctgcc cgagc 45

<210> 41
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 41
ccggaattct gccaaagtatg agccatcctg g 31

<210> 42
<211> 17
<212> DNA
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<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 42
gccatccaga aggtggt 17

<210> 43
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 43

gtcttgtcag ggaagat

17

<210> 44
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 44
ggcaggagac caccttgcga gtgcccac

28

<210> 45
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 45
gggtgggctg taccttctgg aacagggc

28

<210> 46
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 46
ggcgctggct taccggaga ggaatggg

28

<210> 47
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 47
ggaatgggtg ttgtctcctc caaagatgc

28

<210> 48
<211> 1316
<212> DNA
<213> Cricetulus griseus

<400> 48
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<210> 49
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic DNA

<400> 49
 gatcctgctg ggacaaaaat tgg 23

<210> 50
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic DNA

<400> 50
 cttaacatcc caagggatgc tg 22

<210> 51
 <211> 1965
 <212> DNA
 <213> Cricetulus griseus

<400> 51
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aatggccac ttctgatgga aaaaaaaaaa aaaaaaaaaa aaaaaa 1965

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<210> 52

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

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27

<210> 53

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 53

cactgagcca ggggccacac agcatcc

27

<210> 54

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 54

cccctcacgc atgaagcctg gag

23

<210> 55

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 55

tgccaccgtt tcctccataa gcccaga

27

<210> 56

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 56

atggctcaag ctcccgttaa gtgcccga

28

<210> 57

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic DNA

<400> 57

tcaagcgttt gggttgtcc tcatgag

27

<210> 58

<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 58
tccgggatg gcgagatggg caagc 25

<210> 59
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 59
cttgacatgg ctctgggctc caag 24

<210> 60
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 60
ccacttcagt cggtcggtag tattt 25

<210> 61
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 61
cgctcaccg cctgaggcga catg 24

<210> 62
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 62
ggcagggtgct gtcggtgagg tcaccatagt gc 32

<210> 63
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 63

ggggccatgc caaggactat gtcg

24

<210> 64
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic DNA

<400> 64
 atgtggctga tggtacaaaa tgatg

25

<210> 65
 <211> 1504
 <212> DNA
 <213> Cricetulus griseus

<220>
 <221> CDS
 <222> (1).. (1119)

<400> 65
 atg gct cac gct ccc gct agc tgc ccg agc tcc agg aac tct ggg gac 48
 Met Ala His Ala Pro Ala Ser Cys Pro Ser Ser Arg Asn Ser Gly Asp
 1 5 10 15
 ggc gat aag ggc aag ccc agg aag gtg gcg ctc atc acg ggc atc acc 96
 Gly Asp Lys Gly Lys Pro Arg Lys Val Ala Leu Ile Thr Gly Ile Thr
 20 25 30
 ggc cag gat ggc tca tac ttg gca gaa ttc ctg ctg gag aaa gga tac 144
 Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr
 35 40 45
 gag gtt cat gga att gta cgg cga tcc agt tca ttt aat aca ggt cga 192
 Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg
 50 55 60
 att gaa cat tta tat aag aat cca cag gct cat att gaa gga aac atg 240
 Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met
 65 70 75 80
 aag ttg cac tat ggt gac ctc acc gac agc acc tgc cta gta aaa atc 288
 Lys Leu His Tyr Gly Asp Leu Thr Asp Ser Thr Cys Leu Val Lys Ile
 85 90 95 100
 atc aat gaa gtc aaa cct aca gag atc tac aat ctt ggt gcc cag agc 336
 Ile Asn Glu Val Lys Pro Thr Glu Ile Tyr Asn Leu Gly Ala Gln Ser
 105 110 115
 cat gtc aag att tcc ttt gac tta gca gag tac act gca gat gtt gat 384
 His Val Lys Ile Ser Phe Asp Leu Ala Glu Tyr Thr Ala Asp Val Asp
 120 125 130
 gga gtt ggc acc ttg cgg ctt ctg gat gca att aag act tgt ggc ctt 432
 Gly Val Gly Thr Leu Arg Leu Leu Asp Ala Ile Lys Thr Cys Gly Leu
 135 140 145
 ata aat tct gtg aag ttc tac cag gcc tca act agt gaa ctg tat gga 480
 Ile Asn Ser Val Lys Phe Tyr Gln Ala Ser Thr Ser Glu Leu Tyr Gly
 150 155 160
 aaa gtg caa gaa ata ccc cag aaa gag acc acc cct ttc tat cca agg 528
 Lys Val Gln Glu Ile Pro Gln Lys Glu Thr Thr Pro Phe Tyr Pro Arg
 165 170 175 180

tcg ccc tat gga gca gcc aaa ctt tat gcc tat tgg att gta gtg aac 576
 Ser Pro Tyr Gly Ala Ala Lys Leu Tyr Ala Tyr Trp Ile Val Val Asn
 185 190 195

ttt cga gag gct tat aat ctc ttt gcg gtg aac ggc att ctc ttc aat 624
 Phe Arg Glu Ala Tyr Asn Leu Phe Ala Val Asn Gly Ile Leu Phe Asn
 200 205 210

cat gag agt cct aga aga gga gct aat ttt gtt act cga aaa att agc 672
 His Glu Ser Pro Arg Arg Gly Ala Asn Phe Val Thr Arg Lys Ile Ser
 215 220 225

cgg tca gta gct aag att tac ctt gga caa ctg gaa tgt ttc agt ttg 720
 Arg Ser Val Ala Lys Ile Tyr Leu Gly Gln Leu Glu Cys Phe Ser Leu
 230 235 240

gga aat ctg gac gcc aaa cga gac tgg ggc cat gcc aag gac tat gtc 768
 Gly Asn Leu Asp Ala Lys Arg Asp Trp Gly His Ala Lys Asp Tyr Val
 245 250 255 260

gag gct atg tgg ctg atg tta caa aat gat gaa cca gag gac ttt gtc 816
 Glu Ala Met Trp Leu Met Leu Gln Asn Asp Glu Pro Glu Asp Phe Val
 265 270 275

ata gct act ggg gaa gtt cat agt gtc cgt gaa ttt gtt gag aaa tca 864
 Ile Ala Thr Gly Glu Val His Ser Val Arg Glu Phe Val Glu Lys Ser
 280 285 290

ttc atg cac att gga aag acc att gtg tgg gaa gga aag aat gaa aat 912
 Phe Met His Ile Gly Lys Thr Ile Val Trp Glu Gly Lys Asn Glu Asn
 295 300 305

gaa gtg ggc aga tgt aaa gag acc ggc aaa att cat gtg act gtg gat 960
 Glu Val Gly Arg Cys Lys Glu Thr Gly Lys Ile His Val Thr Val Asp
 310 315 320

ctg aaa tac tac cga cca act gaa gtg gac ttc ctg cag gga gac tgc 1008
 Leu Lys Tyr Tyr Arg Pro Thr Glu Val Asp Phe Leu Gln Gly Asp Cys
 325 330 335 340

tcc aag gcg cag cag aaa ctg aac tgg aag ccc cgc gtt gcc ttt gac 1056
 Ser Lys Ala Gln Gln Lys Leu Asn Trp Lys Pro Arg Val Ala Phe Asp
 345 350 355

gag ctg gtg agg gag atg gtg caa gcc gat gtg gag ctc atg aga acc 1104
 Glu Leu Val Arg Glu Met Val Gln Ala Asp Val Glu Leu Met Arg Thr
 360 365 370

aac ccc aac gcc tga gcacctctac aaaaaaattc gcgagacatg gactatggtg 1159
 Asn Pro Asn Ala
 375

cagagccagc caaccagagt ccagccactc ctgagaccat cgaccataaa ccctcgactg 1219
 cctgtgtcgt ccccacagct aagagctggg ccacaggttt gtgggcacca ggacggggac 1279
 actccagagc taaggccact tcgcttttgt caaaggctcc tctcaatgat tttgggaaat 1339
 caagaagttt aaaatcacat actcatttta ctgaaatta tgtcactaga caacttaaat 1399
 ttttgagtct tgagattggt tttctctttt ctatttaaat gatctttcta tgaccagca 1459
 aaaaaaaaaa aaaaaaggga tataaaaaaa aaaaaaaaaa aaaaa 1504

<210> 66
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic DNA

<400> 66
atgaagttgc actatggtga cctca

25

<210> 67
<211> 59
<212> DNA
<213> Cricetulus griseus

<400> 67
ccgacagcac ctgcctagta aaaatcatca atgaagtcaa acctacagag atctacaat 59

<210> 68
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 68
gacttagcag agtacactgc agatg

25

<210> 69
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic DNA

<400> 69
accttgata gaaaggggtg gtctc

25

<210> 70
<211> 125
<212> DNA
<213> Cricetulus griseus

<400> 70
ttgatggagt tggcaccttg cggtctctgg atgcaattaa gacttggtgc cttataaatt 60
ctgtgaagtt ctaccaggcc tcaactagtg aactgtatgg aaaagtgcaa gaaatacccc 120
agaaa 125

<210> 71
<211> 376
<212> PRT
<213> Cricetulus griseus

<400> 71
Met Ala His Ala Pro Ala Ser Cys Pro Ser Ser Arg Asn Ser Gly Asp
1 5 10 15
Gly Asp Lys Gly Lys Pro Arg Lys Val Ala Leu Ile Thr Gly Ile Thr
20 25 30
Gly Gln Asp Gly Ser Tyr Leu Ala Glu Phe Leu Leu Glu Lys Gly Tyr
35 40 45
Glu Val His Gly Ile Val Arg Arg Ser Ser Ser Phe Asn Thr Gly Arg
50 55 60
Ile Glu His Leu Tyr Lys Asn Pro Gln Ala His Ile Glu Gly Asn Met

65	70	75	80
Lys 85	Leu His Tyr Gly 90	Asp Leu Thr Asp Ser Thr 95	Cys Leu Val Lys Ile 100
Ile 105	Asn Glu Val Lys 105	Pro Thr Glu Ile Tyr 110	Asn Leu Gly Ala Gln Ser 115
His 120	Val Lys Ile Ser 120	Phe Asp Leu Ala 125	Glu Tyr Thr Ala Asp Val Asp 130
Gly 135	Val Gly Thr Leu Arg Leu 140	Leu Asp Ala Ile Lys 145	Thr Cys Gly Leu
Ile 150	Asn Ser Val Lys Phe 155	Tyr Gln Ala Ser Thr 160	Ser Glu Leu Tyr Gly
Lys 165	Val Gln Glu Ile Pro 170	Gln Lys Glu Thr Thr 175	Pro Phe Tyr Pro Arg 180
Ser 185	Pro Tyr Gly Ala 185	Ala Lys Leu Tyr Ala 190	Tyr Trp Ile Val Val Asn 195
Phe 200	Arg Glu Ala Tyr Asn Leu Phe 205	Ala Val Asn Gly Ile 210	Leu Phe Asn
His 215	Glu Ser Pro Arg Arg Gly 220	Ala Asn Phe Val Thr 225	Arg Lys Ile Ser
Arg 230	Ser Val Ala Lys Ile Tyr 235	Leu Gly Gln Leu Glu 240	Cys Phe Ser Leu
Gly 245	Asn Leu Asp Ala Lys 250	Arg Asp Trp Gly His 255	Ala Lys Asp Tyr Val 260
Glu 265	Ala Met Trp Leu Met Leu Gln Asn 270	Asp Glu Pro Glu Asp Phe 275	Val
Ile 280	Ala Thr Gly Glu Val His Ser 285	Val Arg Glu Phe Val 290	Glu Lys Ser
Phe 295	Met His Ile Gly Lys Thr 300	Ile Val Trp Glu Gly 305	Lys Asn Glu Asn
Glu 310	Val Gly Arg Cys Lys 315	Glu Thr Gly Lys Ile 320	His Val Thr Val Asp
Leu 325	Lys Tyr Tyr Arg Pro 330	Thr Glu Val Asp Phe 335	Leu Gln Gly Asp Cys 340
Ser 345	Lys Ala Gln Gln Lys Leu Asn Trp 350	Lys Pro Arg Val Ala Phe 355	Asp
Glu 360	Leu Val Arg Glu Met Val Gln 365	Ala Asp Val Glu Leu Met 370	Arg Thr
Asn 375	Pro Asn Ala		

<210> 72
 <211> 321
 <212> PRT
 <213> Cricetulus griseus

<400> 72
 Met Gly Glu Pro Gln Gly Ser Arg Arg Ile Leu Val Thr Gly Gly Ser
 1 5 10 15

Gly Leu Val Gly Arg Ala Ile Gln Lys Val Val Ala Asp Gly Ala Gly
 20 25 30
 Leu Pro Gly Glu Glu Trp Val Phe Val Ser Ser Lys Asp Ala Asp Leu
 35 40 45
 Thr Asp Ala Ala Gln Thr Gln Ala Leu Phe Gln Lys Val Gln Pro Thr
 50 55 60
 His Val Ile His Leu Ala Ala Met Val Gly Gly Leu Phe Arg Asn Ile
 65 70 75 80
 Lys Tyr Asn Leu Asp Phe Trp Arg Lys Asn Val His Ile Asn Asp Asn
 85 90 95
 Val Leu His Ser Ala Phe Glu Val Gly Thr Arg Lys Val Val Ser Cys
 100 105 110
 Leu Ser Thr Cys Ile Phe Pro Asp Lys Thr Thr Tyr Pro Ile Asp Glu
 115 120 125
 Thr Met Ile His Asn Gly Pro Pro His Ser Ser Asn Phe Gly Tyr Ser
 130 135 140
 Tyr Ala Lys Arg Met Ile Asp Val Gln Asn Arg Ala Tyr Phe Gln Gln
 145 150 155 160
 His Gly Cys Thr Phe Thr Ala Val Ile Pro Thr Asn Val Phe Gly Pro
 165 170 175
 His Asp Asn Phe Asn Ile Glu Asp Gly His Val Leu Pro Gly Leu Ile
 180 185 190
 His Lys Val His Leu Ala Lys Ser Asn Gly Ser Ala Leu Thr Val Trp
 195 200 205
 Gly Thr Gly Lys Pro Arg Arg Gln Phe Ile Tyr Ser Leu Asp Leu Ala
 210 215 220
 Arg Leu Phe Ile Trp Val Leu Arg Glu Tyr Asn Glu Val Glu Pro Ile
 225 230 235 240
 Ile Leu Ser Val Gly Glu Glu Asp Glu Val Ser Ile Lys Glu Ala Ala
 245 250 255
 Glu Ala Val Val Glu Ala Met Asp Phe Cys Gly Glu Val Thr Phe Asp
 260 265 270
 Ser Thr Lys Ser Asp Gly Gln Tyr Lys Lys Thr Ala Ser Asn Gly Lys
 275 280 285
 Leu Arg Ala Tyr Leu Pro Asp Phe Arg Phe Thr Pro Phe Lys Gln Ala
 290 295 300
 Val Lys Glu Thr Cys Ala Trp Phe Thr Asp Asn Tyr Glu Gln Ala Arg
 305 310 315 320
 Lys

<210> 73
 <211> 590
 <212> PRT
 <213> Cricetulus griseus

<400> 73
 Met Ala Ser Leu Arg Glu Ala Ser Leu Arg Lys Leu Arg Arg Phe Ser

Ala Glu Leu Gly Leu Gln Ser Ile Ala Phe Ser Val Phe Pro Asn Val
370 375 380
Pro Glu Asp Ser His Glu Lys Pro Cys Val Ile His Ser Ile Leu Asn
385 390 395 400
Ser Gly Cys Cys Val Ala Pro Gly Ser Val Val Glu Tyr Ser Arg Leu
405 410 415
Gly Pro Glu Val Ser Ile Ser Glu Asn Cys Ile Ile Ser Gly Ser Val
420 425 430
Ile Glu Lys Ala Val Leu Pro Pro Cys Ser Phe Val Cys Ser Leu Ser
435 440 445
Val Glu Ile Asn Gly His Leu Glu Tyr Ser Thr Met Val Phe Gly Met
450 455 460
Glu Asp Asn Leu Lys Asn Ser Val Lys Thr Ile Ser Asp Ile Lys Met
465 470 475 480
Leu Gln Phe Phe Gly Val Cys Phe Leu Thr Cys Leu Asp Ile Trp Asn
485 490 495
Leu Lys Ala Met Glu Glu Leu Phe Ser Gly Ser Lys Thr Gln Leu Ser
500 505 510
Leu Trp Thr Ala Arg Ile Phe Pro Val Cys Ser Ser Leu Ser Glu Ser
515 520 525
Val Ala Ala Ser Leu Gly Met Leu Asn Ala Ile Arg Asn His Ser Pro
530 535 540
Phe Ser Leu Ser Asn Phe Lys Leu Leu Ser Ile Gln Glu Met Leu Leu
545 550 555 560
Cys Lys Asp Val Gly Asp Met Leu Ala Tyr Arg Glu Gln Leu Phe Leu
565 570 575
Glu Ile Ser Ser Lys Arg Lys Gln Ser Asp Ser Glu Lys Ser
580 585 590



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